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## WHAT IS CLAIMED IS:

- 1. A method for identifying DNA mutation using microwells which comprises the steps of:
- (i) preparing amplified biotinylated DNA fragments of a portion of nucleotide sequence to be identified by PCR using a biotin-bound primer;
- (ii) preparing a probe comprising normal sequence corresponding to the DNA sequence to be identified;
- (iii) affixing the probe prepared in Step(ii) to the amine group of microwell;
- (iv) adding biotinylated DNA fragments prepared in Step(i) to the probe-affixed microwell;
- (v) adding a streptavidin-linked degradation enzyme to the microwell in order to bind the degradation enzyme to biotin moiety of the probe-captured sample DNA fragment; and,
- (vi) adding a substrate to be reacted with the degradation enzyme and detecting the color or absorbance change caused by degradation of the substrate.
- 2. The method for identifying DNA mutation of claim 1, wherein the probe is a nucleotide sequence consisting of more than 10 nucleotides and containing a phosphate moiety at 5' end.
- 3. The method for identifying DNA mutation of claim 1, wherein Step(iii) comprises adding single-stranded probe to the microwell, adding catalysts in order to bind the phosphate moiety of the single-stranded probe to the amine group of microwell, and washing the microwell.
- 4. The method for identifying DNA mutation of claim 3, wherein the catalysts are ice-cold solutions of 10mM 1-methylimidazole, pH 7.0 and 1-ethyl-3-(3-dimethylaminopropyl)-carbodiimide(EDC), pH 7:0.

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- 5. The method for identifying DNA mutation of claim 3, wherein the washing is performed by employing 0.4M NaOH/0.25% Tween-20 solution.
- 6. The method for identifying DNA mutation of claim l, wherein Step(iv) comprises binding single-stranded DNA fragments obtained in Step(i) to the probe of microwell, removing the residual DNA fragments, and washing the microwell.
  - 7. The method for identifying DNA mutation of claim 6, further comprising pretreatment of the microwell with a solution containing  $dH_2O$ , 20xSSPE/0.0167% Triton X-100 and salmon sperm DNA(10mg/ml) at 50% for 20 minutes before adding the single-stranded DNA fragments obtained in Step(i) to the microwell.
  - 8. The method for identifying DNA mutation of claim 6, wherein binding the DNA fragments obtained in Step(i) to the probe occurs in a solution containing  $dH_2O$ , 20xSSPE/0.0167% Triton X-100 and salmon sperm DNA(10mg/ml).
  - 9. The method for identifying DNA mutation of claim 6, wherein the washing is performed by employing 0.5xSSC/0.1% Tween-20 solution.
    - 10. The method for identifying DNA mutation of claim 1, wherein Step(v) comprises the first washing of the microwell, introducing streptavidin-linked degradation enzyme to the microwell for binding of the enzyme with biotin, removing the residual reaction mixture, and the second washing of the microwell.
- 11. The method for identifying DNA mutation of claim 10, wherein the first washing is performed by employing a buffer solution of 100mM Tris-HCl(pH 7.5) containing 150mM NaCl/0.1% Tween-20.

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- 12. The method for identifying DNA mutation of claim 10, wherein the streptavidin-linked degradation enzyme is streptavidin-alkaline phosphatase dissolved in a buffer solution of 100mM Tris-HCl(pH 7.5) containing 150mM NaCl.
- 13. The method for identifying DNA mutation of claim 10, wherein the second washing comprises treatment of the microwell with a buffer solution of 100mM Tris-HCl(pH 7.5) containing 150mM NaCl/0.1% Tween-20 at  $60\,^{\circ}$ C for 10 minutes.
- 14. The method for identifying DNA mutation of claim 1, wherein the substrate to be reacted with streptavidin-linked degradation enzyme is a synthetic peptide showing color or absorbance change during the degradation.
- 15. The method for identifying DNA mutation of claim 14, wherein the substrate is pNPP(p-nitrophenyl phosphate) provided that the streptavidin-linked degradation enzyme of streptavidin-alkaline phosphatase is employed.
- 16. The method for identifying DNA mutation of claim 1, wherein the color change is detected with naked eyes.
- 25 17. The method for identifying DNA mutation of claim 1, wherein the absorbance is measured by employing an ELISA reader.
- 18. A kit for identifying DNA mutation which 30 comprises:
  - (i) a microwell whose inside has amine group;
  - (ii) 10mM 1-ethyl-3-(3-dimethylaminopropyl)-carbodiimide(EDC), pH 7.0 and 10mM 1-methylimidazole, pH 7.0;
  - (iii) 0.4M NaOH/0.25% Tween-20 solution;
    - (iv) a solution containing  $dH_2O$ , 20xSSPE/0.0167% Triton X-100 and salmon sperm DNA(10mg/ml);

- (v) 0.5xSSC/0.1% Tween-20 solution;
- (vi) streptavidin-alkaline phosphatase;
- (vii) 100 mM Tris-HCl(pH 7.5) solution containing 150 mMNaCl;
- (viii) 100mM Tris-HCl(pH 7.5) solution containing 150mM 5 NaCl/0.1% Tween-20; and,
  - (ix) pNPP(p-nitrophenyl phosphate).